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# Assignment of an essential role for the *Neurospora frequency* gene in circadian entrainment to temperature cycles

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Circadian systems include slave oscillators and central pacemakers, and the cores of eukaryotic circadian clocks described to date are composed of transcription and translation feedback loops (TTFLs). In the model system *Neurospora*, normal circadian rhythmicity requires a TTFL in which a White Collar complex (WCC) activates expression of the *frequency* (*frq*) gene, and the FRQ protein feeds back to attenuate that activation. To further test the centrality of this TTFL to the circadian mechanism in *Neurospora*, we used low-amplitude temperature cycles to compare WT and *frq*-null strains under conditions in which a banding rhythm was elicited. WT cultures were entrained to these temperature cycles. Unlike those normal strains, however, *frq*-null mutants did not truly entrain to the same cycles. Their peaks and troughs always occurred in the cold and warm periods, respectively, strongly suggesting that the rhythm in *Neurospora* lacking *frq* function simply is driven by the temperature cycles. Previous reports suggested that a FRQ-less oscillator (FLO) could be entrained to temperature cycles, rather than being driven, and speculated that the FLO was the underlying circadian-rhythm generator. These inferences appear to derive from the use of a phase reference point affected by both the changing waveform and the phase of the oscillation. Examination of several other phase markers as well as results of additional experimental tests indicate that the FLO is, at best, a slave oscillator to the TTFL, which underlies circadian rhythm generation in *Neurospora*.

FRQ-less oscillator | *frq* | FRQ

Circadian programs in eukaryotes are widely perceived to be the output of multiple oscillatory systems based on cell intrinsic transcription and translation feedback loops (TTFLs) (1–3). In many animals and fungi, heterodimeric PAS domain-containing transcription factors drive expression of genes encoding proteins that block the activity of their heterodimeric activators; such negative feedback loops generally are believed to make up the cores of these circadian clocks. In addition to these autonomous biological clocks, slave oscillators also exist within the panoply of circadian systems. Early studies on entrainment in *Drosophila* gave rise to models in which a pacemaker drove a slave oscillator that directly regulated an overt rhythmic event (4), and noncircadian slaves have since been experimentally described (e.g., ref. 5). However, because there are few molecular descriptions of slave oscillators, their existence and properties have so far chiefly been inferred from the behavior of the circadian system when exposed to Zeitgeber period lengths outside its innate frequency.

At the core of the TTFL in the circadian model system *Neurospora crassa* are the products of the *frequency* (*frq*), *white collar-1* (*wc-1*), and *wc-2* genes. Similar to animal systems, *Neurospora* possesses a feedback loop in which a heterodimeric activator, the White Collar complex (WCC) of the PAS proteins WC-1 and WC-2, activates expression of *frq* and thus FRQ, which in turn depresses transcriptional activation by WCC (6). In this

organism the clock controls several processes including the daily production of asexual spores (conidia). Rhythmic conidiation is visualized in cultures growing along the surface of media as a regularly occurring pattern of aerial hyphae and orange spores (a “band”). In addition to this circadian regulation, the developmental processes leading to conidiation are independently affected by light, temperature, humidity, and media composition, as well as by oscillators that lack full circadian credentials (reviewed in ref. 7). The first such oscillation (in the absence of a fully functioning TTFL) was observed in *frq*-null strains two decades ago (8–10). Observed oscillations in *Neurospora* that lack circadian characteristics and can operate absent the core TTFL have been inferred to be underpinned by operation of FRQ-less oscillators (FLOs) (11, 12), several of which are now believed to exist (reviewed in ref. 7).

Despite these conceptual models, molecular mechanisms of FRQ-less oscillations remain cryptic; the molecular relationship between the driver and the slave has never been clear; and the only sporadic appearance of the original FLO under free-running conditions further rendered its study problematic. To circumvent these obstacles, some studies (e.g., ref. 13) have exploited the observation that, whereas driven rhythms simply respond to environmental cycles, oscillators can entrain to subharmonics of their innate frequencies; that is, a 24-h circadian clock can entrain to a recurring 12-h cycle in the phenomenon known as frequency demultiplication (14). Additionally, temperature cycles have been used to reveal the FLO and to probe the relationship between it and the circadian oscillator (15). This work suggested that the phase of the temperature-induced FLO varied systematically with the period of the temperature cycle and did so in a manner that paralleled the intact circadian system (15). Such results would indicate that this FLO truly can be entrained by temperature cycles. Normal circadian entrainment in *frq*-null strains, i.e., strains lacking the TTFL, would have profound implications. In contemplating these implications, including self-sufficiency for the FLO and that the TTFL might not be necessary for circadian rhythmicity, the authors proposed a novel, ingenious, and speculative model in which the FLO was the underlying circadian oscillator and the TTFL part of the entraining mechanism to light (13, 16). Because this model would require major revisions in circadian theory and in the interpretation of a great many previous results, we reexamined the premise and results on which it was based.

Here, we report the results of experiments performed in several laboratories in the United States and United Kingdom

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Abbreviations: FLO, FRQ-less oscillator; FRP, free running period; TTFL, transcription and translation feedback loop; ZT, Zeitgeber time.

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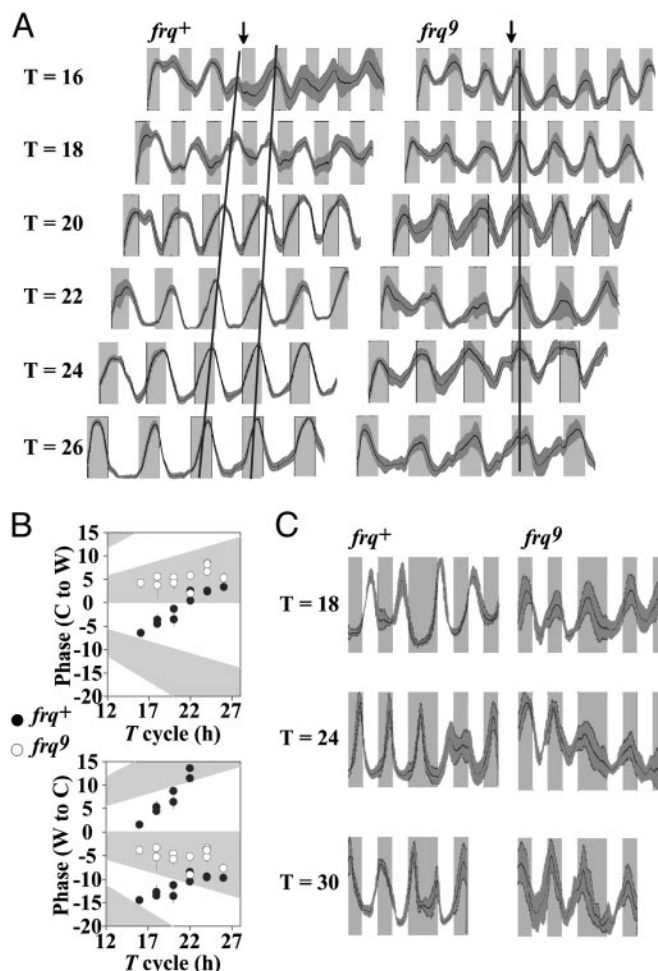




Primary data showing the behavior of clock-mutant and wild-type (WT) strains exposed to low-amplitude temperature cycles are shown in Fig. 1 C–E. In strains bearing functional *frq* alleles, *frq*<sup>2</sup> (FRP ≈ 18 h), *frq*<sup>+</sup> (FRP ≈ 21 h), and *frq*<sup>3</sup> (FRP ≈ 24 h), peaks scan through the warm and cool periods, phase leading or lagging transitions depending on the relationship between the FRP and *T* (Fig. 1 C–E). However, this behavior is not seen in the *frq*-null allele *frq*<sup>0</sup>: Peaks and troughs of conidiation always occur during the cool and warm phases, respectively (see also Fig. 2). These data provide information concerning the nature of the oscillators responding to cycling temperature and suggest that there may be no need to invoke the existence of a FLO underlying the biological cycles observed in *frq*-null strains, even though the banding seen in these strains under temperature cycles shows a *frq*-less oscillation.

It is perhaps valuable to compare these rhythmic phenomena (see also ref. 15) with what occurs in a more typical case in light/dark cycles. In clock WT *Neurospora* cycling in darkness, minimal FRQ levels coincide with the peak in conidiation that normally occurs ≈11 h after the light-to-dark transfer. FRQ levels are high in extended light and fall upon transfer to darkness, with new synthesis of FRQ not observed until a half-circadian cycle later (≈10–14 h after the light-to-dark transition). The FRQ/WCC TTFL responds to shifts in temperature, with FRQ levels increasing rapidly as temperature rises and falling rapidly when the temperature decreases (22). Thus, in a WT strain subjected to 22-h temperature cycles, the shift from cool to warm after 11 h coincides with the FRQ minimum and reinforces the natural momentum of FRQ kinetics; the peak of the conidiation band occurs at the upward temperature transition (Fig. 1D). In longer temperature cycles (*T* > FRP; Fig. 1E), the peak must occur before the transition, because the negative feedback mechanism dictates when FRQ reaches its minimum; conversely, under shorter cycles (*T* < FRP; Fig. 1C) the peak occurs after the transition. For strains with functional circadian clocks but different period lengths, this explanation still holds. For example, in *frq*<sup>2</sup> (FRP ≈ 18 h) the peak occurs at the cool-to-warm transition when the entraining cycle has 9 h at 22°C and 9 h at 27°C (9:9, Fig. 1C) and occurs before the transition in longer cycles (Fig. 1D and E). Absent functional FRQ, temperature cycles still elicit cyclic spore production (Fig. 1 C–E and ref. 15), but in this case spore production appears to respond directly to temperature: Peaks of conidiation always occur in the cool sections of the cycle regardless of cycle length, although the duration of spore production increased with increasing cycle length. Generally, therefore, whether temperature is driving or entraining the clock can be assessed by comparing the phase of the rhythm when exposed to cycles of different period lengths (Fig. 1B).

To accurately assign the function of *frq* in the circadian system, we examined in greater detail the behavior of both the intact (*frq*<sup>+</sup>) circadian system and the FLO within a broad range of temperature cycles (Fig. 2). As before (Fig. 1 and ref. 15), 22°C/27°C temperature cycles were used (Fig. 2A). It is clear that in *frq*<sup>+</sup>, rhythm peaks and troughs scan across the warm and cool periods depending on period length of the entraining cycle; this behavior exemplifies typical entrainment. In the FLO, exposed in a *frq*-null (*frq*<sup>0</sup>) strain, the peak of the rhythm occurs in the cool period, ≈5 h after the warm-to-cool transition, and similarly the trough of the rhythm invariably is found in the midst of the warm period. This invariant behavior indicates that the FLO does not show typical entrainment. All results from WT and *frq*<sup>0</sup> strains from three different laboratories were pooled, and the peaks in the conidiation bands were plotted (Fig. 2B). In this manner of plotting the data, a sloped line reflects a systematic dependence of oscillator phase on the period of the temperature cycle, indicative of entrainment; a flat line reflecting no dependence of phase on *T* indicates a direct response to,



**Fig. 2.** *frq*<sup>+</sup> strains are entrained, and *frq*<sup>0</sup> strains are driven, by temperature cycles. (A) Densitometric tracings of the conidial banding rhythm entrained to 22°C/27°C temperature cycles of varying period length (indicated at left). Shading is as in Fig. 1; middle lines report average (*n* ≥ 6 race tubes) pixel density, and shading above and below this line marks ± 1 SD. The widths of the cool and warm periods were drawn to scale because the growth rate is higher at the higher temperature. The end of the third warm period in each tube has been aligned as indicated by the vertical arrow, and a line was drawn through the third (and in *frq*<sup>+</sup> fourth) peaks to highlight the trends in phase. This line is sloped in WT (Left) showing the systematic change in phase as a function of *T* consistent with entrainment, but the line is vertical in the *frq*-null strain (Right). Similar results were obtained in all three laboratories; plotted data are from one laboratory. (B) The phase of the rhythm peak under different period length (*T*) cycles was measured as the average number of hours the peaks occurred after (–) or before (+) the cool (22°C) to warm (27°C) (Upper) or warm to cool (Lower) transitions for *frq*<sup>0</sup> (○) and *frq*<sup>+</sup> (●). When plotting the phase relative to the warm to cool transition, the *frq*<sup>+</sup> peak cannot always be unambiguously plotted as occurring before or after the transition. Thus, we plotted the results as delays (negative values) as well as advances (positive values). This plot represents all data collected independently in three different laboratories. Each data point is an average phase value from at least three cycles per race tube from *n* ≥ 6 race tubes ± 1 SD. To test how well entrainment period length could predict phase, we performed a linear regression analysis and found a highly significant linear relationship (*P* < 0.001) between phase and *T* in the *frq*<sup>+</sup> strain. However, in *frq*<sup>0</sup> the relationship was not significant (*P* > 0.05), indicating that this strain is not entrained. (C) Densitometric scans of *frq*<sup>+</sup> (Left) and *frq*<sup>0</sup> (Right) strains under cycles of 9 h/9 h (Top), 12 h/12 h (Middle), and 15 h/15 h (Bottom) at 22°C and 27°C. After the fourth full temperature cycle (first two cycles not shown), the temperature was held at 22°C for another half-cycle before resuming regular cycling. In *frq*<sup>+</sup>, the oscillation continues, whereas in *frq*<sup>0</sup>, cycling ceases until the temperature rises again, consistent with the rhythm being driven. Data are from one laboratory, *n* = 6 race tubes ± 1 SD; see also Fig. 5, which is published as supporting information on the PNAS web site.





Table 1. Statistical analysis of data in Fig. 3B

	Slope	SE	Significance*
<i>frq</i> <sup>+</sup>			
Trough	1.70	0.12	$4.9 \times 10^{-19}$
Onset	1.05	0.07	$1.6 \times 10^{-19}$
Peak	0.80	0.06	$6.4 \times 10^{-19}$
Offset	0.91	0.06	$6.8 \times 10^{-19}$
<i>frq</i> <sup>9</sup> and <i>frq</i> <sup>10</sup> combined			
Trough	0.20	0.06	0.001
Onset	0.50	0.07	$1.3 \times 10^{-9}$
Peak	-0.05	0.05	0.33
Offset	0.16	0.07	0.02

Data from all three laboratories (26 separate estimates of phase as a function of genotypes and *T*) were compiled, 11 for *frq*<sup>+</sup> and 15 collectively for *frq*<sup>9</sup> and *frq*<sup>10</sup>. Phase was estimated for each individual race tube from cycling during at least 4 days from a total of 51 race tubes for *frq*<sup>+</sup> and 75 for *frq*<sup>9</sup> and *frq*<sup>10</sup>.

FRP, for instance, cycles that are about half the length of a normal circadian cycle (14, 20). This phenomenon, known as frequency demultiplication, affords an independent evaluation of the oscillator in *frq*-null strains as compared with WT. For instance, when clock WT *Neurospora* is exposed to cycles of 6-h warm/6-h cool, it will exhibit a 24-h rhythm just as if the entraining cycle was 12-h warm/12-h cool (20). We reasoned that this phenomenon would provide another independent test of whether the FLO can exhibit normal entrainment or would instead simply passively respond to the temperature cycles. The period length of the FLO in *frq*<sup>9</sup> is uncompensated, so it depends on temperature and nutrition (8, 9) and can be as short as 12 h (8, 9, 15); alternatively, the *frq*-less oscillations have been modeled as arising from a noise-driven damped harmonic oscillator of 21-h period length (23). We thus used a range of cycle lengths, as short as 6 h (3:3), which would allow entrainment of a 12-h rhythm by demultiplication, and as long as 12 h (6:6), which would demultiply to yield entrainment to 24 h. As expected, clock WT (*frq*<sup>+</sup>) strains successfully entrained by demultiplication. Warm/cool cycles of 4.5/4.5 yielded a rhythm of  $\approx 18$  h, 5/5 cycles led to a rhythm of  $\approx 20$  h, and 6/6 cycles yielded a rhythm of  $\approx 24$  h (Fig. 4); 3/3 cycles could in principle demultiply to a 12-h rhythm, but 12 h is outside of the WT limits of entrainment, so the observed rhythm simply free runs at its normal FRP of  $\approx 21$  h. In contrast, we saw no evidence for frequency demultiplication in the *frq*-null strains to any cycle lengths in this range (see also ref. 13). Instead, at all frequencies, the rhythms observed in *frq*-null strains simply assumed the periodicity of the driving temperature cycle, again showing that strains lacking *frq* are incapable of normal entrainment. A similar loss of demultiplication-driven entrainment has been observed in *per*-null mutants of *Drosophila* (24).

## Discussion

The developmental processes that culminate in the production of aerial hyphae and conidiation occur both in clock-enabled strains of *Neurospora* and strains bearing the null-alleles *frq*<sup>9</sup> or *frq*<sup>10</sup> that lack a circadian core TTFL. In the presence of *frq*-encoded functions, including those specified by any of several (nonnull) missense mutations in this gene (17, 18), conidiation is regulated in a manner bearing all of the hallmarks of a normal circadian rhythm: a robust self-sustained oscillation  $\approx 21$  h in length, with precise control of phase, period, and entrainment to environmental cues, as well as compensation of period against differences in ambient temperature or nutrition. Absent *frq* functions, all of this is lost. Nevertheless, rhythmic output from a *frq*-less oscillator still can sometimes be observed under permissive conditions (8–10). However, the overt rhythm con-

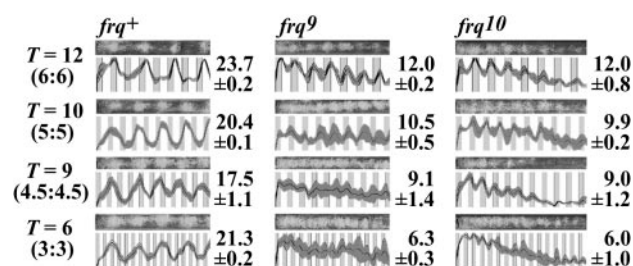


Fig. 4. High-frequency temperature cycles cannot elicit demultiplication in the absence of functional *frq*. *frq*<sup>+</sup> and *frq*-null strains were subjected to temperature cycles (*T*) whose durations ranged downward from the circadian range, as indicated on the far left of the figure. Whereas *frq*<sup>+</sup> is able to demultiply to 18-, 20-, and 24-h periodicities, and free-runs in shorter duration cycles, both *frq*-null strains show conidiation being driven by temperature cycles for all *T*'s applied. On the right of each profile is the period length (in hours)  $\pm$  1 SD estimated from at least six race tubes. Similar experiments yielded equivalent results independently in two laboratories; data reported are from one laboratory.

trolled by such a FLO has lost its robustness, precision of phase, and all aspects of temperature and nutritional compensation; moreover, the period becomes highly variable among consecutive days and shortens with increasing temperature (8–10). However, a rhythm can be dependably visualized by exposing *frq*-null cultures to low-amplitude temperature cycles (15). This process makes it possible to ask whether a bona fide FLO underlies this aspect of *frq*-less rhythmicity by following surrogate markers of it, the various phase reference points. Among these, peaks and troughs are reliable and relevant to the biology of the organism. Onset, however, is not a reliable marker, because it is influenced directly by environmental factors as well as being putatively controlled by a FLO. In any case, by using temperature cycles and reference points of peak and trough as well as offset, we have shown that the FLO is not capable of normal circadian entrainment and, by inference, that the TTFL is required for such entrainment. Given these consistent results from standard reference points, the data suggest that the apparent entrainment seen by using onset is an artifact of the altered waveform. Our results also provide an alternative to “normal entrainment” (13, 15) as the interpretation of the behavior of *frq*-null strains in temperature cycles by showing how use of the onset reference point confounded those phase estimates. Additionally, because we show that the previous results (15) can be duplicated by using the onset reference point, it is unlikely that the discrepancy between our conclusions and those previously proposed lies in any subtle differences in strains, growth conditions, or experimental setup among different laboratories. Finally, we believe that our more extensive data provide more consistent and compelling conclusions.

The data obtained and analyzed in the current study are most consistent with the *frq*-less oscillation being driven by changes in temperature, because all our experiments have failed to find evidence for circadian entrainment of the FLO that was inferred to regulate these biological cycles (15). Absent these data, a role for this FLO as a “circadian rhythm generator,” as previously suggested (13, 16), seems unlikely. An extension of this conclusion is that the FRQ/WCC TTFL is not simply required for light input to a postulated temperature-entrainable oscillator (13) but has a central role in the circadian oscillator.

These considerations leave open the question of the identity of this temperature-influenced *frq*-less oscillation. At present, no known molecular components can be assigned to a FLO (9, 25), so one can only guess about its importance, and even its existence, in WT cells. Without temperature cycles, for instance, oscillations appear in only a fraction of *frq*-null cultures and only

after several days of growth (8, 9, 25). An analogous phenomenon is known in the field of chemical oscillators: When conditions are changed in a mixture capable of robust oscillation, a delayed and sporadic appearance of oscillations often signals the organization of an alternative oscillatory state, one not formed when the regular components are present, which is nucleated after the system has passed an “induction period” (26). From this analogy, Christensen *et al.* (27) have suggested that a FLO may not normally exist and, instead, represents an unnatural oscillatory state formed when normal connections cannot be made. Alternatively, perhaps the FLO is always present but not always overtly manifest. Or it could reflect developmental circuits that normally are organized by the circadian circuitry but, absent a TTFL, can instead be ordered by temperature cycles. It is tempting to liken FLO to an hourglass that can be flipped at an environmental transition; the experiments in Fig. 2C in which FLO ceased to cycle as soon as the temperature cycle disappeared would be consistent with this hypothesis. However, when a FLO does appear during growth in constant conditions, the oscillation generally continues for several days (9, 25). This finding suggests that some oscillator, however weak, must still exist there to restart the cycle again each day.

More generally, it is now known that *Neurospora* harbors a number of FLOs that can appear in the absence of the TTFL

(e.g., refs. 27–29 and reviews in refs. 7 and 18). Although none of these can yet be characterized as wholly circadian, and not any are known to be essential for expression of circadian rhythms, each can display some circadian characteristics. Our data do not preclude a role for other FLOs in the circadian system, but at present a simple and plausible view unifying all of the FLOs is that these are oscillators that can be coupled to a FRQ/WCC-associated circadian core (7). Absent the FRQ/WCC loop, they can run on their own in a noncircadian manner. Whether such FLOs evolved before or after the circadian mechanism, and what their contribution is, if any, to the normal circadian program, are questions that remain unanswered.

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